WEST Search History

DATE: Saturday, January 11, 2003

Set Name side by side		Hit Count	Set Name result set
DB=US	SPT; PLUR=YES; OP=ADJ		
L5	11 and 3-5 exonuclease	0	L5
L4	L3 and (gene silenc\$ or co-suppr\$ or cosuppr\$)	0	L4
L3	3-5 exonuclease	283	L3
L2	ll and plant	53	L2
L1	exonuclease and (gene silenc\$ or co-supp\$ or cosuppr\$)	61	L1

END OF SEARCH HISTORY

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FILE 'HOME' ENTERED AT 15:50:44 ON 11 JAN 2003

=> file agricola caplus biosis
COST IN U.S. DOLLARS

SINCE FILE TOTAL ENTRY SESSION 0.21 0.21

FULL ESTIMATED COST

FILE 'AGRICOLA' ENTERED AT 15:50:53 ON 11 JAN 2003

FILE 'CAPLUS' ENTERED AT 15:50:53 ON 11 JAN 2003 USE IS SUBJECT TO THE TERMS OF YOUR STN CUSTOMER AGREEMENT. PLEASE SEE "HELP USAGETERMS" FOR DETAILS. COPYRIGHT (C) 2003 AMERICAN CHEMICAL SOCIETY (ACS)

FILE 'BIOSIS' ENTERED AT 15:50:53 ON 11 JAN 2003 COPYRIGHT (C) 2003 BIOLOGICAL ABSTRACTS INC.(R)

=> s gene silencing or cosuppression or co-suppression L1 3585 GENE SILENCING OR COSUPPRESSION OR CO-SUPPRESSION

=> s l1 and exonuclease

L2 5 L1 AND EXONUCLEASE

=> dup rem 12 PROCESSING COMPLETED FOR L2

L3 3 DUP REM L2 (2 DUPLICATES REMOVED)

=> d 1-3 ti

- L3 ANSWER 1 OF 3 CAPLUS COPYRIGHT 2003 ACS
- TI cDNA and protein sequences of novel polypeptides comprising a 3'-5' exonuclease domain and methods of controlling gene expression and gene silencing in plants
- L3 ANSWER 2 OF 3 CAPLUS COPYRIGHT 2003 ACS DUPLICATE 1
- TI Molecular characterisation of RecQ homologues in Arabidopsis thaliana
- L3 ANSWER 3 OF 3 CAPLUS COPYRIGHT 2003 ACS DUPLICATE 2
- TI Silencing of .beta.-1,3-glucanase genes in tobacco correlates with an increased abundance of RNA degradation intermediates

=> d pi

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CO, CR, CU, CZ, DE, DK, DM, DZ, EC, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, TZ, UA, UG, US, UZ, VN, YU, ZA, ZW, AM, AZ, BY, KG, KZ, MD, RU, TJ, TM

RW: GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZW, AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, TR, BF, BJ, CF, CG, CI, CM, GA, GN, GQ, GW, ML, MR, NE, SN, TD, TG
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=> d 2 ab

ANSWER 2 OF 3 CAPLUS COPYRIGHT 2003 ACS DUPLICATE 1 L3Members of the RecQ family of DNA helicases are involved in processes AΒ linked to DNA replication, DNA recombination and gene silencing. RecQ homologs of various animals have been described recently. Here, for the first time for plants, we characterized cDNAs of all in all six different RecQ-like proteins that are expressed to different extents in Arabidopsis thaliana. Surprisingly, three of these proteins are small in size [AtRecQl1, AtRecQl2, AtRecQl3-606, 705 and 713 amino acids (aa), resp.], whereas the two bigger proteins result from a duplication event during plant evolution [AtRecQl4A and AtRecQl4B-1150 and 1182 aa, resp.]. Another homolog (AtRecQsim, 858 aa) most probably arose by insertion of an unrelated sequence within its helicase domain. The presence of these homologs demonstrates the conservation of RecQ family functions in higher eukaryotes. We also detected a small gene (AtWRNexo) encoding 285 aa which, being devoid of any RecQ-like helicase domain, reveals a striking homol. to the exonuclease domain of human Werner protein, a prominent RecQ helicase of larger size. By means of the two-hybrid assay we were able to detect an interaction between AtWRNexo and AtRecQl2, indicating that activities that reside in a single protein chain in mammals might in plants be complemented in trans.

=> d 2 so

L3 ANSWER 2 OF 3 CAPLUS COPYRIGHT 2003 ACS DUPLICATE 1 SO Nucleic Acids Research (2000), 28(21), 4275-4282 CODEN: NARHAD; ISSN: 0305-1048

=> d 2 ab

ANSWER 2 OF 3 CAPLUS COPYRIGHT 2003 ACS DUPLICATE 1 L_3 Members of the RecQ family of DNA helicases are involved in processes AB linked to DNA replication, DNA recombination and gene silencing. RecQ homologs of various animals have been described recently. Here, for the first time for plants, we characterized cDNAs of all in all six different RecQ-like proteins that are expressed to different extents in Arabidopsis thaliana. Surprisingly, three of these proteins are small in size [AtRecQl1, AtRecQl2, AtRecQl3-606, 705 and 713 amino acids (aa), resp.], whereas the two bigger proteins result from a duplication event during plant evolution [AtRecQl4A and AtRecQl4B-1150 and 1182 aa, resp.]. Another homolog (AtRecQsim, 858 aa) most probably arose by insertion of an unrelated sequence within its helicase domain. presence of these homologs demonstrates the conservation of RecQ family functions in higher eukaryotes. We also detected a small gene (AtWRNexo) encoding 285 aa which, being devoid of any RecQ-like helicase domain, reveals a striking homol. to the exonuclease domain of human Werner protein, a prominent RecQ helicase of larger size. By means of the two-hybrid assay we were able to detect an interaction between AtWRNexo and AtRecQl2, indicating that activities that reside in a single protein chain in mammals might in plants be complemented in trans.

=> d 2 au

L3 ANSWER 2 OF 3 CAPLUS COPYRIGHT 2003 ACS DUPLICATE 1

AU Hartung, Frank; Plchova, Helena; Puchta, Holger

=> d 3 ab

AB

L3 ANSWER 3 OF 3 CAPLUS COPYRIGHT 2003 ACS DUPLICATE 2

Post-transcriptional gene silencing of .beta.-1,3 qlucanase genes in the transgenic tobacco line T17 is characterized by an increased turnover and, as a consequence, reduced levels of gn1 transgene and endogenous .beta.-1,3 glucanase mRNAs. Here, addnl. gn1 RNAs, both larger and smaller than the full-length messenger, are shown to accumulate in silenced plants of the transgenic tobacco line T17. The longer-than-full-length gn1 RNAs are the result of cryptic processing of the gn1 messenger. The small gn1 RNAs in silenced plants correspond to distal and proximal parts of the mature gn1 messenger. The proximal RNA products are intact at their 5' extremity, but terminate at different positions at the 3'-end. The distal RNA products contain a poly(A) tail and are truncated to various positions at the 5'-end. These observations indicate that degrdn. of the mature gnl transcript does not start at the 5'- or 3'-end, but rather are consistent with degrdn. of the gnl transcript starting with an endonucleolytic cleavage followed by internal exonuclease digestion. Importantly, the truncated products are more abundant in silenced plants than in expressing plants. This suggests, together with the previously reported silencing-related increased gn1 mRNA turnover and the similar rates of gn1 transcription in silenced and expressing T17 plants, that the predominant decay route for the gn1 transcripts differs between silenced and expressing conditions.

=> d 3 so

L3 ANSWER 3 OF 3 CAPLUS COPYRIGHT 2003 ACS DUPLICATE 2

SO Nucleic Acids Research (1998), 26(22), 5176-5181

CODEN: NARHAD; ISSN: 0305-1048

=> s ((levin j?) or (levin, j?))/au L4 1941 ((LEVIN J?) OR (LEVIN, J?))/AU

=> s 14 and exonuclease

L5 6 L4 AND EXONUCLEASE

=> dup rem 15

PROCESSING COMPLETED FOR L5

L6 4 DUP REM L5 (2 DUPLICATES REMOVED)

=> d 1-4 ti

L6 ANSWER 1 OF 4 CAPLUS COPYRIGHT 2003 ACS

TI cDNA and protein sequences of novel polypeptides comprising a 3'-5' exonuclease domain and methods of controlling gene expression and gene silencing in plants

L6 ANSWER 2 OF 4 CAPLUS COPYRIGHT 2003 ACS DUPLICATE 1

TI In vitro detection of endonuclease IV-specific DNA damage formed by bleomycin in vivo

L6 ANSWER 3 OF 4 CAPLUS COPYRIGHT 2003 ACS DUPLICATE 2

TI Analysis of class II (hydrolytic) and class I (.beta.-lyase) apurinic/apyrimidinic endonucleases with a synthetic DNA substrate

- L6 ANSWER 4 OF 4 BIOSIS COPYRIGHT 2003 BIOLOGICAL ABSTRACTS INC.
 TI ENZYMATIC REPAIR OF SPECIFIC OXIDATIVE DAMAGES TO DNA DEOXYRIBOSE IN ESCHERICHIA-COLI.
- => d 2 au
- L6 ANSWER 2 OF 4 CAPLUS COPYRIGHT 2003 ACS DUPLICATE 1
 AU Levin, Joshua D.; Demple, Bruce
- => d 2 ab
- ANSWER 2 OF 4 CAPLUS COPYRIGHT 2003 ACS L6 DUPLICATE 1 Endonuclease IV of Escherichia coli has been implicated by genetic studies AB in the repair of DNA damage caused by the antitumor drug bleomycin, but the lesion(s) recognized by this enzyme in vivo have not been identified. We used the sensitive primer activation assay, which monitors the formation of 3'-OH groups that support in vitro synthesis by E. coli DNA polymerase I, to det. whether endonuclease IV-specific damage could be detected in the chromosomal DNA of cells lacking the enzyme after in vivo treatment with bleomycin. Chromosomal DNA isolated after a 1 h bleomycin treatment from wild-type, endonuclease IV-deficient (nfo-) and endonuclease IV-overproducing (p-nfo .apprx.10-fold) strains all supported modest polymerase activity. However, in vitro treatment with purified endonuclease IV activated subsequent DNA synthesis with samples from the nfo- strain (an av. of 2.6-fold), to a lesser extent for samples from wild-type cells (2.1-fold), and still less for the p-nfo samples (1.5-fold). This pattern is consistent with the presence of unrepaired damage that correlates inversely with the in vivo activity of endonuclease IV. Incubation of the DNA from bleomycin-treated nfo- cells with polymerase and dideoxynucleoside triphosphates lowered the endonuclease IV-independent priming activity, but did not affect the amt. of activation seen after endonuclease IV treatment. Primer activation with DNA from the nfo- strain could also be obtained with purified E. coli exonuclease III in vitro, but a quant. comparison demonstrated that endonuclease IV was .gtoreq.5-fold more active in this assay. Thus, endonuclease IV-specific damage can be detected after in vivo exposure to bleomycin. These may be 2-deoxypentos-4-ulose residues, but other possibilities are discussed.

=> d ti

L7 ANSWER 1 OF 1 CAPLUS COPYRIGHT 2003 ACS
TI cDNA and protein sequences of novel polypeptides comprising a 3'-5'
exonuclease domain and methods of controlling gene expression and
gene silencing in plants

=> s ((phillips k?) or (phillips, k?)/au
UNMATCHED LEFT PARENTHESIS '((PHILLIPS'
The number of right parentheses in a query must be equal to the number of left parentheses.

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=> s ((phillips k?) or (phillips, k?))/au
L8 1007 ((PHILLIPS K?) OR (PHILLIPS, K?))/AU
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=> s 18 and exonuclease L9 1 L8 AND EXONUCLEASE

=> d ti ANSWER 1 OF 1 CAPLUS COPYRIGHT 2003 ACS 1.9 cDNA and protein sequences of novel polypeptides comprising a 3'-5' TΤ exonuclease domain and methods of controlling gene expression and gene silencing in plants => s ((glazov e?) or ((glazov, e?))/au UNMATCHED LEFT PARENTHESIS '((GLAZOV' The number of right parentheses in a query must be equal to the number of left parentheses. => s ((glazov e?) or (glazov, e?))/au 23 ((GLAZOV E?) OR (GLAZOV, E?))/AU => s l10 and exonuclease 1 L10 AND EXONUCLEASE 1.11 => d ti L11 ANSWER 1 OF 1 CAPLUS COPYRIGHT 2003 ACS cDNA and protein sequences of novel polypeptides comprising a 3'-5' exonuclease domain and methods of controlling gene expression and gene silencing in plants => s 110 and (gene silen? or cosupp? or co-supp?) 4 L10 AND (GENE SILEN? OR COSUPP? OR CO-SUPP?) => dup rem 12 PROCESSING COMPLETED FOR L2 3 DUP REM L2 (2 DUPLICATES REMOVED) => d 1-3 ti L13 ANSWER 1 OF 3 CAPLUS COPYRIGHT 2003 ACS cDNA and protein sequences of novel polypeptides comprising a 3'-5' exonuclease domain and methods of controlling gene expression and gene silencing in plants L13 ANSWER 2 OF 3 CAPLUS COPYRIGHT 2003 ACS DUPLICATE 1 Molecular characterisation of RecQ homologues in Arabidopsis thaliana L13 ANSWER 3 OF 3 CAPLUS COPYRIGHT 2003 ACS DUPLICATE 2 Silencing of .beta.-1,3-glucanase genes in tobacco correlates with an increased abundance of RNA degradation intermediates => d 2 au L13 ANSWER 2 OF 3 CAPLUS COPYRIGHT 2003 ACS DUPLICATE 1 Hartung, Frank; Plchova, Helena; Puchta, Holger => d 2 kwic L13 ANSWER 2 OF 3 CAPLUS COPYRIGHT 2003 ACS DUPLICATE 1 Members of the RecQ family of DNA helicases are involved in processes linked to DNA replication, DNA recombination and gene silencing. RecQ homologs of various animals have been described recently. Here, for the first time for plants, we characterized cDNAs of.

. . small gene (AtWRNexo) encoding 285 aa which, being devoid of any

RecQ-like helicase domain, reveals a striking homol. to the **exonuclease** domain of human Werner protein, a prominent RecQ helicase of larger size. By means of the two-hybrid assay we were.

- ST RecQl1 RecQl2 RecQl3 RecQl4 DNA helicase Arabidopsis cDNA sequence; exonuclease WRNexo Arabidopsis cDNA sequence
- IT Molecular association

(two-hybrid anal. was able to detect interactions between the WRNexo exonuclease and the RecQl2 helicase)

IT 37228-74-3, Exonuclease

RL: BSU (Biological study, unclassified); PRP (Properties); BIOL (Biological study)

(cloning, sequence and characterization of RecQ homologs in Arabidopsis thaliana)

=> s plant and transgenic and exonuclease L14 9 PLANT AND TRANSGENIC AND EXONUCLEASE

=> dup rem 114
PROCESSING COMPLETED FOR L14
L15 8 DUP REM L14 (1 DUPL)

8 DUP REM L14 (1 DUPLICATE REMOVED)

=> d 1-8 tui

'TUI' IS NOT A VALID FORMAT

In a multifile environment, a format can only be used if it is valid in at least one of the files. Refer to file specific help messages or the STNGUIDE file for information on formats available in individual files.

REENTER DISPLAY FORMAT FOR ALL FILES (FILEDEFAULT):ti

- L15 ANSWER 1 OF 8 CAPLUS COPYRIGHT 2003 ACS
- TI Genes essential for early growth of Arabidopsis thaliana and their use in the development of novel herbicides
- L15 ANSWER 2 OF 8 CAPLUS COPYRIGHT 2003 ACS
- TI Whole cell engineering by mutagenizing a substantial portion of a starting genome and combining mutations with optional reiteration
- L15 ANSWER 3 OF 8 CAPLUS COPYRIGHT 2003 ACS
- TI Detection of nucleic acids by selective depolymerization of probes hybridized to a target sequence and detection of specific hydrolysis products
- L15 ANSWER 4 OF 8 CAPLUS COPYRIGHT 2003 ACS
- TI Generation of genetic vaccines and immunomodulatory polynucleotides by non-stochastic directed evolution techniques
- L15 ANSWER 5 OF 8 BIOSIS COPYRIGHT 2003 BIOLOGICAL ABSTRACTS INC.
- TI Interaction between composite elements in the napA promoter: Both the B-box ABA-responsive complex and the RY/G complex are necessary for seed-specific expression.
- L15 ANSWER 6 OF 8 AGRICOLA DUPLICATE 1
 - TI Interaction between composite elements in the napA promoter: both the B-box ABA-responsive complex and the RY/G complex are necessary for seed-specific expression.
 - L15 ANSWER 7 OF 8 CAPLUS COPYRIGHT 2003 ACS
 - TI Silencing of :beta.-1,3-glucanase genes in tobacco correlates with an increased abundance of RNA degradation intermediates
 - L15 ANSWER 8 OF 8 CAPLUS COPYRIGHT 2003 ACS
 - TI Application of PCR to transgenic plants

=> d 7 so

L15 ANSWER 7 OF 8 CAPLUS COPYRIGHT 2003 ACS SO Nucleic Acids Research (1998), 26(22), 5176-5181 CODEN: NARHAD; ISSN: 0305-1048

=> d 8 ab

L15 ANSWER 8 OF 8 CAPLUS COPYRIGHT 2003 ACS

AB Advantages and disadvantages of PCR are discussed in consideration of plant systems and in comparison to Northern and Southern anal., genomic sequencing, chem. mutagenesis, nucleotide insertions, and exonuclease digestions.

=> d 6 ab

L15 ANSWER 6 OF 8 AGRICOLA DUPLICATE 1 During seed maturation, the transcriptional activity of napin genes is regulated by developmental signals involving the transcriptional activator ABI3 and abscisic acid (ABA). To localize cis elements involved in the seed-specific activity of the napin napA promoter, a systematic analysis was performed focusing on two major element complexes, the B-box and RY/G. Substitution mutation analysis using promoter-reporter gene fusions in stable transgenic tobacco showed synergistic interactions between elements within these complexes. The distal part of the B-box shows similarities to abscisic acid response elements and the proximal portion contains a CA-rich element. In vitro studies involving Exonuclease III protection and electrophoretic mobility shift assays revealed binding by nuclear proteins to elements within the B-box. The distal and proximal parts of the B-box were found to bind distinct nuclear protein complexes. By gain-of-function analysis with a tetramer of the B-box fused to a truncated (-46) cauliflower mosaic virus (CaMV) 35S minimal promoter, it was demonstrated that the B-box mediates strong activity in seeds. Further, it was shown that the elements in the B-box constitute an ABA-responsive complex, since the B-box tetramer mediates ABA-responsiveness in vegetative tissues to a construct containing the CaMV virus 35S enhancer (-343 to -90). Thus, the seed-specific activity of the napA promoter relies on the combinatorial interaction between the RY/G complex and the B-box ABA-responsive complex during the ABA response in seed development.

=> s 3-5 (adj5) exonuclease and plant MISSING OPERATOR '3-5 (ADJ5' The search profile that was entered contains terms or nested terms that are not separated by a logical operator.

=> s 3-5 (adj5) exonuclease MISSING OPERATOR '3-5 (ADJ5' The search profile that was entered contains terms or nested terms that are not separated by a logical operator.

=> dis his

(FILE 'HOME' ENTERED AT 15:50:44 ON 11 JAN 2003)

FILE 'AGRICOLA, CAPLUS, BIOSIS' ENTERED AT 15:50:53 ON 11 JAN 2003
L1 3585 S GENE SILENCING OR COSUPPRESSION OR CO-SUPPRESSION
L2 5 S L1 AND EXONUCLEASE
L3 3 DUP REM L2 (2 DUPLICATES REMOVED)
L4 1941 S ((LEVIN J?) OR (LEVIN, J?))/AU

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6 S L4 AND EXONUCLEASE
L5
              4 DUP REM L5 (2 DUPLICATES REMOVED)
L6
L7
              1 S L6 AND (GENE SILEN? OR CO-SUP? OR COSUP?)
L8
          1007 S ((PHILLIPS K?) OR (PHILLIPS, K?))/AU
              1 S L8 AND EXONUCLEASE
L9
             23 S ((GLAZOV E?) OR (GLAZOV, E?))/AU
L10
L11
              1 S L10 AND EXONUCLEASE
              4 S L10 AND (GENE SILEN? OR COSUPP? OR CO-SUPP?)
L12
              3 DUP REM L2 (2 DUPLICATES REMOVED)
L13
              9 S PLANT AND TRANSGENIC AND EXONUCLEASE
L14
              8 DUP REM L14 (1 DUPLICATE REMOVED)
L15
=> s l1 and plant
         1274 L1 AND PLANT
=> s ll and review
          405 L1 AND REVIEW
=> d 1-10 ti
L17 ANSWER 1 OF 405 AGRICOLA
     The efficacy of RNAi in the study of the plant cytoskeleton.
    ANSWER 2 OF 405 AGRICOLA
     New advances in understanding the molecular biology of plant/potyvirus
TТ
     interactions.
    ANSWER 3 OF 405 AGRICOLA
L17
     A model for RNA-mediated gene silencing in higher
     plants.
L17 ANSWER 4 OF 405 AGRICOLA
     Can we explain RNA-mediated virus resistance by homology-dependent
     gene silencing?
L17 ANSWER 5 OF 405 AGRICOLA
```

- The silence of genes in transgenic plants.
- L17 ANSWER 6 OF 405 AGRICOLA
- Molecular data and the dynamic nature of polyploidy.
- L17 ANSWER 7 OF 405 AGRICOLA
- Genomic imprinting in plants: parental effects and trans-inactivation TI phenomena.
- L17 ANSWER 8 OF 405 AGRICOLA
- Mating-type gene switching in Saccharomyces cerevisiae. ΤI
- L17 ANSWER 9 OF 405 CAPLUS COPYRIGHT 2003 ACS
- RNA interference of HIV replication TI
- L17 ANSWER 10 OF 405 CAPLUS COPYRIGHT 2003 ACS
- TT Gene Silencing-Based Disease Resistance
- => d 3 so
- L17 ANSWER 3 OF 405 AGRICOLA
- Plant molecular biology, May 1998. Vol. 37, No. 2. p. 349-362 Publisher: Dordrecht: Kluwer Academic Publishers. CODEN: PMBIDB; ISSN: 0167-4412

L17 ANSWER 1 OF 405 AGRICOLA

SO Journal of plant growth regulation, Dec 2000. Vol. 19, No. 4. p. 371-384 Publisher: New York: Springer-Verlag New York, c1982-CODEN: JPGRDI; ISSN: 0721-7595

=> d ab

L17 ANSWER 1 OF 405 AGRICOLA

AB Recent studies on a variety of organisms point to the ubiquity of RNA interference (RNAi) as a means to induce a gene-specific block to translation. RNAi has gained popularity in the last few years in the study of a number of problems in development. In this review, we highlight recent findings with RNAi using several different kinds of animals and fungi, and we show how these responses parallel cosuppression effects described in plants nearly a decade earlier. We then point to the efficacy of RNAi in studying minor and regulatory components of the plant cytoskeleton, and we highlight some recent studies using this approach with the water fern, Marsilea vestita.